



PCT

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/509,738

DATE: 10/12/2004

TIME: 12:09:20

Input Set : A:\pto.lm.txt

Output Set: N:\CRF4\10122004\J509738.raw

3 <110> APPLICANT: Director-General of National Institute of Advanced Industrial Science and Technology; Info Genes Co., Ltd.; Kazusa DNA Research Institute
 W--> 4 nce and Technology; Info Genes Co., Ltd.; Kazusa DNA Research Institute
 W--> 5 <120> TITLE OF INVENTION: Application of KIAA0172 gene functions for therapeutics, diagnosis,
 W--> 6 and pharmaceuticals
 W--> 7 <130> FILE REFERENCE: PH-1610-PCT
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/509,738
 C--> 10 <141> CURRENT FILING DATE: 2004-09-30
 12 <150> PRIOR APPLICATION NUMBER: JP 2002/99422
 13 <151> PRIOR FILING DATE: 2002-04-01
 15 <160> NUMBER OF SEQ ID NOS: 49
 17 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

Does Not Comply
 Corrected Diskette Needed
 (PS, 1-2)

792 <210> SEQ ID NO: 49
 793 <211> LENGTH: 24
 794 <212> TYPE: DNA
 795 <213> ORGANISM: Artificial Sequence
 797 <220> FEATURE:
 798 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic DNA
 800 <400> SEQUENCE: 49
 801 cctaaacct ctataatata caac
 E--> 802 1/27

24

BEST AVAILABLE COPY

VERIFICATION SUMMARY

DATE: 10/12/2004

PATENT APPLICATION: US/10/509,738

TIME: 12:09:21

Input Set : A:\pto.lm.txt

Output Set: N:\CRF4\10122004\J509738.raw

L:5 M:283 W: Missing Blank Line separator, <120> field identifier
L:7 M:283 W: Missing Blank Line separator, <130> field identifier
L:9 M:270 C: Current Application Number differs, Replaced Application Number
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:295 M:283 W: Missing Blank Line separator, <220> field identifier
L:379 M:283 W: Missing Blank Line separator, <400> field identifier
L:493 M:283 W: Missing Blank Line separator, <220> field identifier
L:579 M:283 W: Missing Blank Line separator, <400> field identifier
L:633 M:283 W: Missing Blank Line separator, <220> field identifier
L:690 M:283 W: Missing Blank Line separator, <220> field identifier
L:776 M:283 W: Missing Blank Line separator, <400> field identifier
L:802 M:254 E: No. of Bases conflict, LENGTH:Input:27 Counted:25 SEQ:49
L:802 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:802 M:252 E: No. of Seq. differs, <211> LENGTH:Input:24 Found:25 SEQ:49